

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/551,341

CRF Processing Date: 2/4/2002
 Edited by: 16
 Verified by: (STIC staff)

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FEB 06 2002

TECH CENTER 1600/2900

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



1646

RAW SEQUENCE LISTING

DATE: 02/04/2002

PATENT APPLICATION: US/09/551,341

TIME: 18:41:42

Input Set : N:\Crf3\01242002\I551341.raw

Output Set: N:\CRF3\02042002\I551341.raw

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SEQUENCE LISTING

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1 (1) GENERAL INFORMATION:
2   (i) APPLICANT: Brass, Clifford A., Glue, Paul W., and Piken,
3       Edward.
4   (ii) TITLE OF INVENTION: HCV Combination Therapy
5   (iii) NUMBER OF SEQUENCES: 4
6   (iv) CORRESPONDENCE ADDRESS:
7       (A) ADDRESSEE: Schering-Plough Corporation
8       (B) STREET: 2000 Galloping Hill Road
9       (C) CITY: Kenilworth
10      (D) STATE: New Jersey
11      (E) COUNTRY: USA
12      (F) ZIP: 07033
13   (v) COMPUTER READABLE FORM:
14       (A) MEDIUM TYPE: Floppy disk
15       (B) COMPUTER: Macintosh
16       (C) OPERATING SYSTEM: 7.5.3
17   (vi) CURRENT APPLICATION DATA:
C--> 18       (A) APPLICATION NUMBER: US/09/551,341
C--> 19       (B) FILING DATE: 18-Apr-2000
20       (C) CLASSIFICATION:
21   (viii) ATTORNEY/AGENT INFORMATION:
22       (A) NAME: Hoffman, Thomas D.
23       (B) REGISTRATION NUMBER: 28,221
24       (C) REFERENCE/DOCKET NUMBER: IN01023K
25   (ix) TELECOMMUNICATION INFORMATION:
26       (A) TELEPHONE: 908-298-5037
27       (B) TELEFAX: 908-298-5388
28 (2) INFORMATION FOR SEQ ID NO: 1:
29   (i) SEQUENCE CHARACTERISTICS:
30       (A) LENGTH: 22 base pairs
31       (B) TYPE: nucleic acid
32       (C) STRANDEDNESS: single
33       (D) TOPOLOGY: linear
W--> 34   (ii) MOLECULE TYPE: DNA (oligonucleotide)
35   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
36       GTGGTCTGCG GAACCGGTGA GT 22
38 (2) INFORMATION FOR SEQ ID NO: 2:
39   (i) SEQUENCE CHARACTERISTICS:
40       (A) LENGTH: 20 base pairs
41       (B) TYPE: nucleic acid
42       (C) STRANDEDNESS: single
43       (D) TOPOLOGY: linear

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RAW SEQUENCE LISTING

DATE: 02/04/2002

PATENT APPLICATION: US/09/551,341

TIME: 18:41:42

Input Set : N:\Crf3\01242002\I551341.raw

Output Set: N:\CRF3\02042002\I551341.raw

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W--> 44      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      45      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
      46      TGCACGGTCT ACGAGACCTC 20
      48 (2) INFORMATION FOR SEQ ID NO: 3:
      49      (i) SEQUENCE CHARACTERISTICS:
      50          (A) LENGTH: 21 base pairs
      51          (B) TYPE: nucleic acid
      52          (C) STRANDEDNESS: single
      53          (D) TOPOLOGY: linear
W--> 54      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      55      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
      56      CTGTGAGGAA CTACTGTCTT C 21
      58 (2) INFORMATION FOR SEQ ID NO: 4:
      59      (i) SEQUENCE CHARACTERISTICS:
      60          (A) LENGTH: 21 base pairs
      61          (B) TYPE: nucleic acid
      62          (C) STRANDEDNESS: single
      63          (D) TOPOLOGY: linear
W--> 64      (ii) MOLECULE TYPE: DNA (oligonucleotide)
      65      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
      66      CCCTATCAGG CAGTACCACA A 21
```

VERIFICATION SUMMARY

DATE: 02/04/2002

PATENT APPLICATION: US/09/551,341

TIME: 18:41:43

Input Set : N:\Crf3\01242002\I551341.raw

Output Set: N:\CRF3\02042002\I551341.raw

L:18 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:19 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:34 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:44 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:54 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:64 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4